Relationship between Parkinson's disease and Cardio-cerebrovascular diseases: a Mendelian randomized study

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Supplement II

Statistical Interpretation:

Inverse variance weighted (IVW): IVW is MR's method for summarizing effect values at multiple loci when analyzing multiple SNPs. Its arithmetic formula is as follows:

The causal effect values calculated for each genetic variant k for the exposure factor and the outcome variable are expressed as the following ratios:

 Y_k / X_k

For the total causal effect values of all instrumental variables, they were assessed by the fixed effect model in the meta-analysis with the following equation:

$$\hat{\beta}_{IVW} = \frac{\sum_{k} X_{k} Y_{k} \sigma_{Yk}^{-2}}{\sum_{k} X_{k}^{2} \sigma_{Yk}^{-2}}$$

The corresponding standard errors are calculated as follows:

$$se(\hat{\beta}_{IVW}) = \sqrt{\frac{1}{\sum_{k} X_{k}^{2} \sigma_{Yk}^{-2}}}$$

MR-Egger: MR-Egger is based on the principle of building a linear regression model with the effect value of the genetic variable as the independent variable and the relationship between the exposure factor and the outcome variable as the dependent variable. The causal effects were estimated by the Egger regression method and tested for the presence of horizontal bias. Its arithmetic formula is as follows:

$$X_{i} = \sum_{j=1}^{J} \gamma_{j} G_{ij} + U_{i} + \varepsilon_{i}^{X}$$
$$Y_{i} = \sum_{j=1}^{J} \alpha_{j} G_{ij} + \beta X_{i} + U_{i} + \varepsilon_{i}^{X}$$

$$\frac{\sum_{j=1}^{J} \hat{\gamma}_{j}^{2} \sigma_{Y_{j}}^{-2} \hat{\beta}_{j}}{\sum_{j=1}^{J} \hat{\gamma}_{j}^{2} \sigma_{Y_{j}}^{-2}}$$
$$\beta + \frac{\sum_{j=1}^{J} \hat{\gamma}_{j} \sigma_{Y_{j}}^{-2} \alpha_{j}}{\sum_{j=1}^{J} \hat{\gamma}_{j}^{2} \sigma_{Y_{j}}^{-2}} = \beta + Bias(\alpha, \gamma)$$

Weighted median: The effect estimates for each SNP were ranked and the weighted median was calculated as the estimate of the causal effect. Its arithmetic formula is as follows:

$$w'_{j} = \frac{\hat{\gamma}_{j}^{2}}{\sigma_{Y_{j}}^{2}}$$

The simple model: Selecting a strongly associated SNP as an instrumental variable and using the effect size of this SNP to estimate the causal effect of exposure on outcome. Its arithmetic formula is as follows:

$$\hat{\beta}_{simple} = \frac{\hat{\beta}_{outcome}}{\hat{\beta}_{exp\,osure}}$$

The weighted mode: The weighted mode is a method for combining the results of multiple Mendelian randomization estimates, which weights the causal effects of different genetic variants on a trait and then takes the weighted plurality as the final causal effect estimate. Its arithmetic formula is as follows:

$$\hat{\beta}_{j} = \frac{\hat{\Gamma}_{j}}{\hat{\gamma}_{j}}$$

Cochran's Q test: Cochran's Q test is a heterogeneity test that is used to test for significant differences between three or more groups of frequencies or rates. Its arithmetic formula is as follows:

$$Q = \sum_{j} Q_{j} = \sum_{j} w'_{j} (\hat{\beta}_{j} - \hat{\beta})^{2}$$

F-statistics: F-statistics is a common statistic in MR Studies, which is generally used to test whether the regression results are significant. Its arithmetic formula is as

follows:

 $F = R^2 \times (N - k - 1) / [(1 - R^2) \times k]$

 R^2 is calculated as $R^2 = 2 \times beta 2 \times (1 - EAF)$ EAF, where beta 2 is an estimate of the genetic effect of each SNP on PD, and EAF is the frequency of the effect allele.

References

- Bowden J, Davey Smith G, Haycock PC, Burgess S. Consistent Estimation in Mendelian Randomization with Some Invalid Instruments Using a Weighted Median Estimator. Genet Epidemiol. 2016 May;40(4):304-14. doi: 10.1002/gepi.21965. Epub 2016 Apr 7. PMID: 27061298; PMCID: PMC4849733.
- Bowden J, Davey Smith G, Burgess S. Mendelian randomization with invalid instruments: effect estimation and bias detection through Egger regression. Int J Epidemiol. 2015 Apr;44(2):512-25. doi: 10.1093/ije/dyv080. Epub 2015 Jun 6. PMID: 26050253; PMCID: PMC4469799.